

Table SI. Analysis of linkage disequilibrium between four SNPs of *FMO3*.

SNP	c.855C>T			c.441C>T			c.923A>G		
	r ²	D'	P-value	r ²	D'	P-value	r ²	D'	P-value
c.441C>T	0.2237	0.8010	<0.001	-	-	-	-	-	-
c.923A>G	0.2565	0.8577	<0.001	0.0632	-0.8301	<0.001	-	-	-
c.472G>A	0.3239	0.8492	<0.001	0.0842	-0.8442	<0.001	0.5332	0.8289	<0.001

SNP, single nucleotide polymorphism; D' = $D/D_{\max}$, D, the coefficient of LD (-0.25 to 0.25); -, not applicable.